

420 Rec'd PCT/PTO 26 JAN 2000

## SEQUENCE LISTING

<110> The Administrators of the Tulane Educational Fund  
Landry, Samuel J.

<120> PREDICTION, DETECTION AND DESIGN OF  
T-CELL EPITOPE

<130> 07005/003WO2

<140> NOT KNOWN

<141> 1998-07-28

<150> 60/054,156

<151> 1997-07-29

<160> 2

<170> FastSEQ for Windows Version 3.0

<210> 1.

<211> 513

<212> PRT

<213> Bacteriophage T4

<400> 1

Met Arg Val Lys Gly Ile Arg Lys Asn Tyr Gln His Leu Trp Lys Trp  
1 5 10 15  
Gly Ile Met Leu Leu Gly Ile Leu Met Ile Cys Ser Ala Ala Glu Asn  
20 25 30  
Leu Trp Val Thr Val Tyr Tyr Gly Val Pro Val Trp Lys Glu Ala Thr  
35 40 45  
Thr Thr Leu Phe Cys Ala Ser Asp Ala Lys Ala Tyr Asp Thr Glu Val  
50 55 60  
His Asn Val Trp Ala Thr His Ala Cys Val Pro Thr Asp Pro Asn Pro  
65 70 75 80  
Gln Glu Val Val Leu Glu Asn Val Thr Glu Asn Phe Asn Met Trp Lys  
85 90 95  
Asn Asn Met Val Glu Gln Met His Glu Asp Ile Ile Ser Leu Trp Asp  
100 105 110  
Gln Ser Leu Lys Pro Cys Val Lys Leu Thr Pro Leu Cys Val Thr Leu  
115 120 125  
Asn Cys Thr Asn Val Asn Val Thr Asn Leu Lys Asn Glu Thr Asn Thr  
130 135 140  
Asn Ser Ser Ser Gly Gly Glu Lys Met Glu Glu Gly Glu Met Lys Asn  
145 150 155 160  
Cys Ser Phe Asn Val Thr Thr Leu Ile Arg Asn Lys Arg Lys Thr Glu  
165 170 175  
Tyr Ala Leu Phe Tyr Lys Leu Asp Val Met Pro Ile Asp His Asp Asn  
180 185 190  
Thr Ser Tyr Thr Leu Ile Asn Cys Lys Ser Ser Thr Ile Thr Gln Ala  
195 200 205

Cys Pro Lys Val Ser Phe Glu Pro Ile Pro Ile His Tyr Cys Ala Pro  
 210 215 220  
 Ala Gly Phe Ala Ile Leu Lys Cys Asn Asp Lys Lys Phe Asn Gly Lys  
 225 230 235 240  
 Gly Pro Cys Lys Asn Val Ser Thr Val Gln Cys Thr His Gly Ile Arg  
 245 250 255  
 Pro Val Val Ser Thr Gln Leu Leu Leu Asn Gly Ser Leu Ala Glu Glu  
 260 265 270  
 Asp Ile Val Ile Arg Ser Glu Asn Phe Thr Asp Asn Ala Lys Asn Ile  
 275 280 285  
 Ile Val Gln Leu Asn Val Ser Ile Glu Ile Asn Cys Thr Arg Pro Asn  
 290 295 300  
 Asn Asn Thr Arg Lys Lys Ile Thr Leu Gly Pro Gly Arg Val Leu Tyr  
 305 310 315 320  
 Thr Thr Gly Glu Ile Ile Gly Asp Ile Arg Arg Ala His Cys Asn Leu  
 325 330 335  
 Ser Arg Thr Ser Trp Asn Asn Thr Leu Lys Gln Ile Val Glu Lys Leu  
 340 345 350  
 Arg Glu Ile Lys Gln Phe Lys Asn Lys Thr Ile Val Phe Lys Gln Ser  
 355 360 365  
 Ser Gly Gly Asp Pro Glu Ile Val Met His Ser Phe Asn Cys Gly Gly  
 370 375 380  
 Glu Phe Phe Tyr Cys Asn Ser Thr Gln Leu Phe Asn Ser Thr Trp His  
 385 390 395 400  
 Ala Asn Gly Thr Trp Lys Asn Thr Glu Gly Ala Asp Asn Asn Ile Thr  
 405 410 415  
 Leu Pro Cys Arg Ile Lys Gln Ile Ile Asn Arg Trp Gln Glu Val Gly  
 420 425 430  
 Lys Ala Met Tyr Ala Pro Pro Ile Glu Gly Gln Ile Arg Cys Leu Ser  
 435 440 445  
 Asn Ile Thr Gly Leu Leu Leu Thr Arg Asp Gly Gly Ser Ser Glu Glu  
 450 455 460  
 Asn Gln Thr Glu Ile Phe Arg Pro Gly Gly Asn Met Lys Asp Asn  
 465 470 475 480  
 Trp Arg Ser Glu Leu Tyr Lys Tyr Lys Val Val Lys Ile Glu Pro Leu  
 485 490 495  
 Gly Val Ala Pro Thr Lys Ala Lys Arg Arg Val Val Gln Arg Glu Lys  
 500 505 510  
 Arg

<210> 2  
 <211> 112  
 <212> PRT  
 <213> Human Immunodeficiency Virus

<400> 2  
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 Ile Leu Val Ser Glu Pro Ala Gln Ala Gly Asp Glu Glu Val Thr Glu  
 20 25 30  
 Ser Gly Leu Ile Ile Gly Lys Arg Val Gln Gly Glu Val Pro Glu Leu  
 35 40 45

Cys Val Val His Ser Val Gly Pro Asp Val Pro Glu Gly Phe Cys Glu  
50 55 60  
Val Gly Asp Leu Thr Ser Leu Pro Val Gly Gln Ile Arg Asn Val Pro  
65 70 75 80  
His Pro Phe Val Ala Leu Gly Leu Lys Gln Pro Lys Glu Ile Arg Lys  
85 90 95  
Gln Lys Phe Val Thr Cys His Tyr Lys Ala Ile Pro Cys Leu Tyr Lys  
100 105 110